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Cryopreservation and Long-Term Storage of Human Low Density Lipoproteins

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Summary: A simple and effective technique of long-term storage of human low density lipoproteins (LDL) has been developed. The technique involves the addition of 1.4 mol/l (or 10% by volume) dimethyl sulphoxide directly to a solution of the freshly isolated LDL in high salt buffer, and subsequent freezing and storage for up to 2 years at -70°C . We have shown that freshly isolated LDL, "preserved" as described above, are able to keep their native properties for a long period, i. e.:

- a) electrophoretic behaviour in non-denaturing (or with sodium dodecyl sulphate, 1 g/l) polyacrylamide 2–16% gradient gel electrophoresis;
- b) immunoreactivity of apolipoprotein B (analyzed by radial immunodiffusion, electroimmunoassay and immunoturbidimetric assay);
- c) immunogeneity of apolipoprotein B;
- d) an average size of LDL particles (analyzed by electron microscopy);
- e) ability to bind with B₂E-receptors of human skin fibroblasts.

The technique can also be applied to radiolabelled LDL samples. Taking into consideration the labour- and time-consuming procedure of obtaining and characterizing LDL, and the preferred use of single well-characterized LDL preparation, we recommend that the above technique of LDL long-term storage be applied in various clinical and biomedical studies.

Introduction

Low density lipoproteins (LDL) play a key role in the transport of cholesterol to the peripheral tissues where they are bound to cellular receptors and ingested by endocytosis (1). LDL also play an important role in the pathological uptake and deposition of cholesterol since very high concentrations of LDL are implicated as the causative agent of some forms of human atherosclerosis, while moderate elevations over long periods of time may be an important factor in the development of most human atherosclerosis (2, 3). Therefore, LDL themselves and their B₂E-receptor binding ability are widely studied (4–8). Furthermore, purified LDL preparations

are also used to obtain poly- and monoclonal antibodies (9–10), as well as a primary standard for apolipoprotein B measurements by immunochemical assays (11). Traditionally, only fresh LDL are used for these purposes. It necessitates a large number of repeated technical operations to obtain and characterize LDL (particularly, when LDL are used as an immunogen and as a primary standard). Hence, an investigator is not able to compare within one long-term experiment the properties of LDL obtained by different methods and time.

Recently, it has been shown that sucrose can be used as a cryoprotector for a long-term storage of the isolated LDL at low temperature (12). On the other hand, there

exist widely known techniques of human and animal cells long-term storage by freezing them in the presence of liquid cryoprotectors (dimethyl sulphoxide, glycerol and etc.) and keeping them at -70°C or, preferably, in liquid nitrogen (13–14).

In this work we suggest a simple and effective technique of long-term LDL storage at -70°C by freezing them in the presence of 1.4 mol/l (10% by volume) dimethyl sulphoxide. It is shown that this storage method does not cause the loss of the native properties of LDL by using them as an immunogen for obtaining the polyclonal monospecific sheep's antiserum, as a primary standard for apolipoprotein B measurements by immunochemical assays and as a ligand. The data reported here have been obtained as a result of two-year experiments.

Materials and Methods

Reagents

Phenyl methylsulphonyl fluoride, sodium bromide, and dimethyl sulphoxide were obtained from Merck (Darmstadt, Germany). Carrier-free Na^{125}I was obtained from Amersham (Arlington Heights, IL). Reagents for polyacrylamide gel electrophoresis were from Serva (Heidelberg, Germany). Horseradish peroxidase (EC 1.11.1.7) and peroxidase substrate (3,3'-diaminobenzidine) were obtained from Sigma Chemical Company (St. Louis, MO).

The candidate international standard plasma pool for apolipoprotein quantitation No. 1883 was obtained from the Centers for Disease Control (Atlanta, Georgia). A reference serum (Lot No. 063511) with assigned values for apolipoprotein A-I (1.62 g/l) and apolipoprotein B (1.21 g/l) was obtained from Behringwerke AG (Marburg, Germany).

Collection of specimens

Blood was collected from hyperlipidaemic and normolipidaemic individuals by venipuncture into Vacutainer tubes (Becton-Dickinson, Rutherford, NJ) in the absence of ethylenediamine-tetraacetate, following a 12–14 h fast, and pooled. Upon separation at 4°C by low-speed centrifugation, the serum was prepared for the immediate isolation of low density lipoprotein fraction. Disodium ethylenediamine-tetraacetate, sodium azide, and phenyl methylsulphonyl fluoride were added to final concentration of 1 mmol/l, 0.2 g/l, and 1 mmol/l, respectively.

Isolation and characterization of LDL

LDL (d 1.030–1.050 kg/l) was isolated from pooled serum of individuals by sequential ultracentrifugation as previously described (15) in a L8-70 ultracentrifuge (Beckman Instruments, Palo Alto, U.S.A.) using 45Ti rotor with potassium bromide being substituted for sodium bromide. Immediately after recentrifugation at d 1.050 kg/l, the LDL fraction was filtered through a filter stack of glassfibre prefilter and 0.2 micron filter (Sartorius GmbH, Göttingen, Germany) under sterile conditions. LDL purity was assessed by 1 g/l sodium dodecyl sulfate, 2–16% gradient polyacrylamide gel electrophoresis (16) and by *Ouchterlony's* method (17) with Calbiochem Corp. (San Diego, CA) anti-apolipoprotein A-I, anti-apolipoprotein A-II and anti-apolipoprotein E antibodies, and with home-made anti-apolipoprotein B antibodies.

Total LDL protein concentration was determined by the method of Lowry et al. (18) as modified by Albers et al. (19). A standard curve was prepared by diluting the bovine plasma albumin standard II (1.37 g/l, Lot No. 34522, Bio-Rad Laboratories, Richmond, CA) to 0.046, 0.091, 0.137, and 0.228 g/l in 0.15 mol/l sodium chloride. Samples were diluted with 0.15 mol/l sodium chloride to three different concentrations within the working range.

Total cholesterol, unesterified cholesterol and triacylglycerols were measured by enzymatic methods using Boehringer (Mannheim, Germany) reagents. Phospholipids were determined as phosphorus (20) using an average relative molecular mass of $M_r = 775$ for phospholipid.

LDL used in receptor-binding studies was radiolabelled with ^{125}I using the iodine monochloride procedure of McFarlane (21) as modified by Bilheimer et al. (22). The specific radioactivity of the apolipoprotein B-100 was about 400 counts/min \cdot ng of protein, and over 97% of the radioactivity was precipitable by incubation with 100 g/l trichloroacetic acid solution at 4°C .

Lipoprotein-deficient serum was obtained by double ultracentrifugation of the freshly pooled serum of healthy donors with a density of 1.25 kg/l for 48 h at 105 000 g. After centrifugation, the serum was dialyzed against 0.15 mol/l sodium chloride, sterilized by filtration (Sartorius filters, 0.2 μm), frozen, and stored at -40°C . The concentration of apolipoprotein B-containing lipoproteins in the lipoprotein-deficient serum did not exceed 0.1 mg/l.

Preparation of antiserum and antibody-enzyme conjugate

Isolated LDL was used to prepare anti-apolipoprotein B antiserum by immunization in sheeps as previously described (23). The specificity of the antisera obtained was assessed by double radial immunodiffusion and two-dimensional immunoelectrophoresis with home-made apolipoprotein A-I, apolipoprotein A-II, apolipoprotein E, apolipoprotein C's, apolipoprotein E, apolipoprotein(a), and with lipoprotein-deficient serum. The IgG fraction for Western blotting was obtained by precipitation with saturated ammonium sulphate as described (24). Rabbit anti-sheep IgG antibodies were conjugated with horseradish peroxidase as previously described (24).

Cryopreservation and storage of LDL

Each LDL preparation was divided into 16–18 portions (0.3 ml). 33 μl of dimethyl sulphoxide per 0.3 ml (10% by volume) was added to the first 12–14 portions and 33 μl of sodium chloride/sodium bromide solution with the density 1.05 kg/l per 0.3 ml (10% by volume) was added to the remaining four portions which were used as controls. Both dimethyl sulphoxide and sodium chloride/sodium bromide solution with the density 1.05 kg/l were previously filtered through a filter stack of glassfibre prefilter and 0.2 micron filter (Sartorius GmbH, Göttingen, Germany) under sterile conditions. Two from four control portions were stored at 4°C . Two other control portions and the portions of the LDL with dimethyl sulphoxide were placed into a foam plastic container with a wall thickness of 2 cm and slowly frozen (about 1°C per min) in the ultra low freezer (Sanyo, Japan) at -70°C as recommended for preservation of human or animal cells (13, 14). Immediately before the use, the LDL portions were quickly defrosted at 37°C .

Polyacrylamide gradient gel electrophoresis

The apolipoprotein B fragmentation was studied on home-made slab 2–16% polyacrylamide linear gradient gels containing 1 g/l sodium dodecyl sulphate (16). The samples were heated for 5 min at 100°C in 50 mmol/l Tris, pH 6.8, 30 g/l sodium dodecyl sulphate, 200 g/l glycerol, 50 g/l 2-mercaptoethanol. The electrophoresis was carried out at 20 mA for 20 h. The proteins were either stained with Coomassie Blue R-250 or transferred to nitrocellulose paper (see below).

Non-denaturing gradient polyacrylamide gel electrophoresis of LDL was done on Pharmacia Electrophoresis Apparatus (GE 2/4) using home-made slab 2–16% polyacrylamide linear gradient gels in 14 mmol/l Tris, 110 mmol/l glycine, 0.1 g/l sodium azide, pH 8.3, as described by *Krauss & Burke* (25). The gels were prerun 30 min at 50 V. Routinely, the samples containing 15 µg of protein were applied to each sample well. Electrophoresis was started at 30 V for 1 h and continued for 20 h at 80 V. Gels were stained with Coomassie Blue R-250 for protein. In some cases LDL samples were prestained with acetylated Sudan Black B. Unstained gels were used for immunoblotting techniques.

Immunoblotting

After electrophoresis, the LDL particles were blotted (26) to nitrocellulose paper (0.45 µm, Serva, Heidelberg, Germany) by the electrophoretic transfer of the gel in the electrophoretic buffer at 100 mA for up to 72 h at 4 °C. All additional operations were performed as described (27).

Immunochemical assays

Single radial immunodiffusion was performed with commercially available M-Partigen-Apolipoprotein B plates from Behringwerke AG (Marburg, Germany) according to the manufacturer's instructions for maximal diffusion time of 96 h at 37 °C. Electroimmunoassay was performed by the method of *Laurell* (28) using 10 g/l agarose (Sigma Chemical Company, St. Louis, MO), containing 15 g/l antiserum to human apolipoprotein B. The LDL samples were diluted to three different concentrations within the working range. All subsequent operations were performed as described (29). Apolipoprotein B values in the LDL samples were also determined by the immunoturbidimetric assay as described (30) with some modifications. Briefly, the LDL samples were diluted to three different concentrations with phosphate-buffered saline (per litre: 9 g of sodium chloride, 10 mmol of monopotassium phosphate, pH 7.4) within the working range (0.3–2.5 g/l). Antiserum was diluted 10-fold in phosphate-buffered saline containing 40 g of polyethylene glycol 6000 (Serva, Heidelberg, Germany) per litre and preincubated for 30 min at 20–25 °C. Then, 25 µl of diluted LDL was added to 750 µl of diluted antiserum. Samples were vortexed, incubated at 20–25 °C for 60 min, vortexed again and the changes in turbidity of the samples against a reagent blank (mixture of 25 µl of sample diluent and 750 µl of diluted antiserum) were measured at 340 nm using a Spectronic 2000 spectrophotometer (Bausch & Lomb, Rochester, New York). A standard curve generated by least-squares linear regression analysis was used to calculate the concentrations of the unknowns.

Standards, quality controls and samples were analyzed in triplicate in all immunochemical, protein, and lipid assays.

Cells

Human skin fibroblasts (6th–10th passages) were grown from forearm skin biopsies of healthy donors. The cells were cultured at 37 °C in an atmosphere of 5% CO₂/95% air in plastic flasks with *Eagles's* minimum essential medium containing 25 mmol/l Hepes, 220 mmol/l sodium bicarbonate, 10 g/l non-essential amino acids, 100 mg/l canamycin, 2 mmol/l glutamine, and 150 g/l fetal calf serum (medium A). The cells were then plated onto 60-mm Petri dishes at a cell density of 1×10^5 cells per dish. Three or four days later, the cells were washed with phosphate-buffered saline and the medium of the same composition as medium A but containing 100 g/l lipoprotein-deficient serum instead of fetal calf serum (medium B) was added and cells were incubated for 48 h.

Measurements of LDL binding, internalization and degradation

The amount of surface-bound ¹²⁵I-labelled LDL, and intracellular ¹²⁵I-labelled LDL was measured in intact fibroblast monolayers

essentially as described by *Goldstein, Basu & Brown* (8). Typically, after incubating the cells at 37 °C for 3 h in the presence of various amounts of ¹²⁵I-labelled LDL (5, 10 or 20 mg/l of [¹²⁵I]LDL protein) for the specific binding, and ¹²⁵I-labelled LDL and a 20-fold excess of unlabelled LDL for the non-specific binding, cell monolayers were washed at 4 °C 5 times with phosphate-buffered saline containing 2 g/l bovine serum albumin and twice with buffer without bovine serum albumin. The cells were then incubated for 1 h with 2 ml of the medium B, containing dextran sulphate (10 g/l), at 4 °C with shaking. The amount of LDL bound was determined from the dextran sulphate-releasable radioactivity in the medium. The cells were then washed and dissolved in 100 mmol/l sodium hydroxide for determination of LDL internalization and protein content. LDL degradation was determined as the difference between the amount of non-iodine, non-lipid trichloroacetic acid-soluble radioactivity in the medium and the amount due to spontaneous degradation. The latter was measured in duplicate incubations with no cells present in the system.

Electron microscopy

Samples of LDL were negatively stained with 20 g/l sodium phosphotungstate (pH 7.4) and immediately examined in the electron microscope as previously described (31).

Results

Electrophoretic investigations

The data of non-denaturing 2–16% gradient polyacrylamide gel electrophoresis show that by the storage of freshly isolated LDL ("fresh" LDL) within 40–60 days at 4 °C particles were formed, having less electrophoretic mobility, well stained by Sudan Black, Coomassie Blue R-250 and interacting with sheep anti-human apolipoprotein B antibodies by immunostaining (see fig. 1, A–C, lane 2). The samples of the same LDL, containing 1.4 mol/l dimethyl sulphoxide and freeze-dried as described in Materials and Methods ("preserved" LDL) after 2 years of storage at –70 °C did not contain such particles (see fig. 1, A–C, lane 3) and had an electrophoretic pattern similar to freshly isolated LDL (see fig. 1, A–C, lane 1). The freezing and subsequent storage at –70 °C within 2 years of the same LDL not containing dimethyl sulphoxide ("control" LDL) led to the formation of particles migrating significantly slower than LDL (see fig. 1, A–C, lane 4) and having similar abilities to be stained by Sudan Black, Coomassie Blue R-250 and interact with sheep anti-human apolipoprotein B antibodies as compared with particles formed under long-term storage of LDL at 4 °C. It is obvious that the particles formed in both cases are aggregated LDL.

To examine probable apolipoprotein B fragmentation we have analyzed the samples of freshly isolated LDL, "fresh", "preserved" and "control" LDL (the latter two – after two-year storage at –70 °C) by 2–16% gradient polyacrylamide gel electrophoresis in the presence of 1 g/l sodium dodecyl sulphate with the subsequent immunoblottings as described in Materials and Methods.

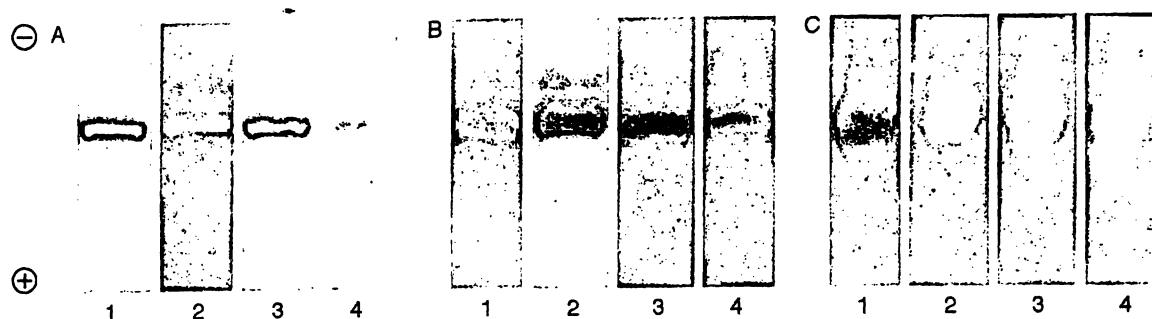


Fig. 1 Non-denaturing 2–16% gradient polyacrylamide gel electrophoresis of: freshly isolated LDL (A–C, line 1), “fresh” LDL – stored for 2 months at 4 °C (A–C, lane 2), “preserved” LDL – frozen in the presence of 1.4 mol/l dimethyl sulfoxide and subsequently stored for 2 years at –70 °C (A–C, lane 3), “control” LDL – frozen without dimethyl sulfoxide and subsequently stored for 2 years at –70 °C (A–C, lane 4). Twenty µg of LDL

were electrophoresed in all cases. Gels stained for lipoprotein with Sudan Black (A), for protein with Coomassie Blue R-250 (B) or LDL samples were immunodetected on a blot of gel by the sheep polyclonal anti-human-apolipoprotein B immunoglobulins and then by horseradish peroxidase-conjugated rabbit anti-sheep IgG antibodies added followed by colour development (C).

The data obtained show no significant differences between the analyzed samples and no apolipoprotein B fragmentation (data not shown).

Similar results were obtained for three different LDL preparations. Hence the given data prove that the freezing of freshly isolated native LDL in the presence of 1.4 mol/l dimethyl sulfoxide and their subsequent storage at –70 °C for two-year period did not cause any significant changes in electrophoretic pattern.

In addition, it has been demonstrated by non-denaturing 2–16% gradient polyacrylamide gel electrophoresis that freezing and subsequent thawing of the “preserved” LDL according to the procedure described in Materials and Methods did not cause changes in electrophoretic pattern even after three-fold repetition. In contrast, the freeze-thawing of “control” LDL (not containing dimethyl sulfoxide) led to the formation of aggregated LDL, whereas the aggregation degree was highly dependent on the number of freeze-thawing cycles. The gels were stained by Sudan Black; similar results were obtained by staining by Coomassie Blue R-250 (data not shown). Thus, the given data indicate a certain antiaggregatory effect of dimethyl sulfoxide on LDL even after three-fold repetition of freeze-thawing procedure.

To study the effect of freezing and subsequent storage in the presence of dimethyl sulfoxide upon the electrophoretic pattern of radiolabelled LDL, the fresh LDL sample was radiolabelled with ^{125}I as described in Materials and Methods. One portion was subsequently frozen in the presence of 1.4 mol/l dimethyl sulfoxide and stored at –70 °C within 7 days (“preserved” [^{125}I]LDL), whereas another one was stored at 4 °C (“freshly isolated” [^{125}I]LDL). Equal aliquots of both “freshly isolated” and “preserved” [^{125}I]LDL were analyzed by non-denaturing 2–16% gradient polyacrylamide gel electrophoresis as described in Materials and Methods. After

electrophoresis, the LDL particles were stained by Coomassie R-250 or transferred to nitrocellulose paper as described in Materials and Methods. After drying, autoradiography was performed at 4 °C using an intensifying screen for 10 min. No differences were found in the electrophoretic patterns for both “freshly isolated” and “preserved” [^{125}I]LDL samples.

Lipid and protein composition

To study the impact of freezing and subsequent storage in the presence of dimethyl sulfoxide upon the LDL lipid and protein content, we compared the results (for three various LDL preparations – LDL1, LDL2, LDL3) of the total and free cholesterol, phospholipid, triacylglycerol and protein determinations, obtained immediately after LDL isolation (“freshly isolated” LDL), and after freezing with 1.4 mol/l dimethyl sulfoxide as described in Materials and Methods, and subsequent storage for 2 years at –70 °C (“preserved” LDL). Table 1 shows that there is no significant difference between the three cases.

Thus, freezing LDL in the presence of dimethyl sulfoxide and subsequent long-term storage at –70 °C have no effect on the results of the LDL lipid and protein determinations.

Immunogeneity of LDL

The LDL samples frozen in the presence of dimethyl sulfoxide (“preserved” LDL) after various periods (from 1 month up to 2 years) of storage at –70 °C were used as an antigen to obtain polyclonal sheep antisera against apolipoprotein B. The analysis of specificity by double radial immunodiffusion and two-dimension immunoelectrophoresis shows that there is no difference between the above antisera and the one obtained by

Tab. 1 Effect of LDL freezing with 1.4 mol/l dimethyl sulphoxide and subsequent storage at -70°C on lipid and protein quantitation^a.

Analyte	Concentration of analyte, g/l					
	LDL1		LDL2		LDL3	
	freshly isolated	preserved	freshly isolated	preserved	freshly isolated	preserved
Total cholesterol	38.3 ± 0.4	38.1 ± 0.8	22.6 ± 0.5	22.8 ± 0.4	34.8 ± 0.4	35.0 ± 0.5
Free cholesterol	10.8 ± 0.3	10.9 ± 0.3	6.3 ± 0.3	6.4 ± 0.3	10.7 ± 0.4	10.8 ± 0.4
Triacylglycerols	6.1 ± 0.2	6.2 ± 0.2	3.9 ± 0.1	3.8 ± 0.1	5.9 ± 0.2	6.0 ± 0.2
Phospholipid	22.3 ± 0.6	22.6 ± 0.7	14.6 ± 0.5	14.4 ± 0.4	23.8 ± 0.6	23.6 ± 0.6
Protein	23.5 ± 0.7	23.7 ± 0.5	13.4 ± 0.4	13.6 ± 0.4	22.0 ± 0.5	22.1 ± 0.5

^a Three LDL preparations (LDL1, LDL2 and LDL3) were obtained as described in Materials and Methods. Aliquots of LDL were assessed for lipid and protein content immediately after isolation ("freshly isolated") and after freezing and subsequent

storage in the presence of 1.4 mol/l dimethyl sulphoxide for 2 years at -70°C ("preserved"). All values are means \pm SD from triplicate determinations.

freshly isolated LDL. The titre determination by an immunoturbidimetric assay (approximately 0.5 g/l) also showed no significant differences between the antisera. We later successfully used the antisera obtained with "preserved" LDL in kits for immunoturbidimetric apolipoprotein B determination in human sera by mass screening.

Hence the LDL freezing in the presence of dimethyl sulphoxide and subsequent long-term storage at -70°C do not exert significant effects upon LDL immunogenic properties.

Apolipoprotein B immunoreactivity

To study the influence of the freezing and storage procedures upon the apolipoprotein B immunoreactivity, we have performed apolipoprotein B analysis of two different LDL preparations (LDL1 and LDL2) by three immunoprecipitin methods: single radial immunodiffusion, electroimmunoassay and immunoturbidimetric assay. LDL samples were analyzed immediately after isolation ("freshly isolated" LDL), after freezing and storage for up to 2 years at -70°C both in the presence of 1.4 mol/l dimethyl sulphoxide ("preserved" LDL) and in the absence of dimethyl sulphoxide ("control" LDL). As shown in table 2 the results of the apolipoprotein B determination by three analytical methods in "freshly isolated" LDL and in "preserved" LDL did not differ significantly (at $p > 0.01$) between both LDL preparations (LDL1 and LDL2). In contrast, the apolipoprotein B values in "control" LDL determined by "gel" techniques (single radial immunodiffusion and electroimmunoassay) significantly differ (at $p < 0.01$) from the ones for "freshly isolated" and "preserved" LDL. Depending upon the method (single radial immunodiffusion or electroimmunoassay) the difference ranged from 16–

23%. A similar difference was not observed for immunoturbidimetric assay (see tab. 2).

Thus, freezing with 1.4 mol/l dimethyl sulphoxide and subsequent storage of LDL for up to 2 years at -70°C do not effect the apolipoprotein B immunoreactivity, at least in the three immunoassays used.

Cell studies

To study the effect of LDL freezing with 1.4 mol/l dimethyl sulphoxide and subsequent storage upon the LDL-receptor activity, equal aliquots of "freshly isolated" and "preserved" [^{125}I]LDL were used in the experiments to measure the specific binding, internalization and degradation of LDL by fibroblasts as described in Materials and Methods. As shown in table 3 the values of specific binding, internalization and degradation of LDL by fibroblasts did not significantly (at $p > 0.01$) differ for "freshly isolated" and "preserved" [^{125}I]LDL, thus indicating the absence of a considerable effect of the [^{125}I]LDL freezing and storage in the presence of 1.4 mol/l dimethyl sulphoxide on LDL-receptor activity.

Electron microscopy

To study the effect of freezing and storage in the presence of dimethyl sulphoxide upon the LDL morphologic properties, one of the two aliquots of LDL was examined in the electron microscope within 1 week after isolation, and the second one after freezing with 1.4 mol/l dimethyl sulphoxide as described in Materials and Methods, and subsequent storage for 2 years at -70°C (data not shown). The particles in both samples appeared to be relatively uniform in size. In both cases the average size of particles was 22 ± 2 nm, which was in

Tab. 2 Effect of LDL freezing with 1.4 mol/l dimethyl sulphoxide and subsequent storage at -70°C on apolipoprotein B quantitation by immunoprecipitin methods^a.

Method	Concentration of apolipoprotein B, g/l					
	LDL1			LDL2		
	freshly isolated	preserved	control	freshly isolated	preserved	control
Radial immunodiffusion	23.4 ± 0.8	23.6 ± 0.6	19.7 ± 0.7	13.7 ± 0.5	13.4 ± 0.5	11.3 ± 0.6
Electroimmunoassay	23.8 ± 0.7	24.2 ± 1.0	18.4 ± 0.8	13.7 ± 0.4	13.4 ± 0.4	10.4 ± 0.3
Immunoturbidimetric assay	23.6 ± 1.0	23.5 ± 0.8	23.2 ± 0.8	13.8 ± 0.3	13.9 ± 0.3	13.9 ± 0.3

^a Two LDL preparations (LDL1 and LDL2) were obtained as described in Materials and Methods. Aliquots of LDL were assessed for apolipoprotein B content immediately after isolation ("freshly isolated") and after freezing and subsequent storage for 2 years at -70°C in the presence ("preserved") or in the absence ("control") of 1.4 mol/l dimethyl sulphoxide. All values are means \pm SD from triplicate determinations.

Tab. 3 Comparison of "freshly isolated" and "preserved" [^{125}I]LDL (freezing with 1.4 mol/l dimethyl sulphoxide and subsequent storage for 7 days at -70°C) specific binding by human skin fibroblasts^a, internalization and degradation at 37°C .

Lipoprotein	ng of [^{125}I]-labelled lipoprotein per mg cell protein		
	Binding	Internalization	Degradation
Freshly isolated [^{125}I]LDL			
10 mg/l	75.0 ± 1.2	679.8 ± 8.3	1026.4 ± 28.5
20 mg/l	149.7 ± 3.1	1109.5 ± 26.2	1734.8 ± 36.3
Preserved [^{125}I]LDL			
10 mg/l	76.9 ± 2.1	683.4 ± 9.8	1013.7 ± 25.6
20 mg/l	146.9 ± 2.3	1124.8 ± 22.1	1716.6 ± 33.7

^a Human fibroblasts were incubated for 3 h at 37°C with the indicated concentrations of radiolabelled lipoproteins. The binding, internalization and degradation were determined as described in Materials and Methods and were corrected for non-specific bind-

ing, internalization and degradation in the presence of 20-fold excess of unlabelled human LDL. All values are means \pm SD from triplicate determinations.

agreement with reported data on particle sizes of such LDL subfraction (31).

Thus, the LDL freezing in the presence of 1.4 mol/l dimethyl sulphoxide and their subsequent storage for up to 2 years at -70°C do not lead to an obvious alteration of morphologic LDL properties.

Discussion

In a majority of research and clinical studies which involved LDL isolation, the researchers have always stressed the necessity to use only freshly isolated LDL, due to the rather quick alterations occurring in the LDL native properties by storage. Thus, for example, to study various immunochemical LDL properties (32), including LDL application as a primary standard for apolipoprotein B measurements by immunochemical assays (11, 19, 23), to study LDL-receptor activity (33, 34), physical and chemical (35, 36) as well as other LDL properties it is usually recommended to conduct the experiments using LDL up to 2–4 weeks after their isolation

and storage at 4°C in the presence of various preservatives. Taking into consideration the labour- and time-consuming process of LDL isolation and characterization as well as the frequent necessity of using one well-characterized portion of LDL, for numerous scientific experiments (for example, in long-term comparative studies) we conclude that the task of long-term LDL storage is a topical and important one.

Aggregation, oxidizing degradation, proteolysis and lipolysis can be major causes for the alterations of native LDL properties occurring during long-term storage. It is obvious that conventional freezing can dramatically decrease the contribution of the latter three processes to LDL denaturation. Nevertheless, according to our electrophoretic data, irreversible formation of aggregated LDL particles occurs upon freezing without an evident apolipoprotein B fragmentation in these particles in contrast to oxidizing (with Cu^{2+}) LDL degradation (37), where, as the authors believe, the generated aggregates present LDL particles cross-linked via apolipoprotein B fragments. The formation of LDL aggregates

was observed by their intensive vortexing as well (38). It is very likely that the apparent loss of apolipoprotein B immunoreactivity we observed after long-term storage of LDL at 4 °C (for more than 1 month) and similar losses reported previously (39, 40) after LDL freezing and subsequent storage at -20 °C can be explained by LDL aggregation. It should also be noted that such apparent losses of apolipoprotein B immunoreactivity were observed only by using "gel" techniques to determine apolipoprotein B: single radial immunodiffusion (39, 40), and single radial immunodiffusion and electroimmunoassay in the present study. This conclusion is supported by the known dependence of the "rocket" size upon the size heterogeneity of albumin and haptoglobin (monomers or dimers) when analyzing them by electroimmunoassay (41), as well as by the recently reported data showing the impact of the LDL particles size upon the divergent values between the results of apolipoprotein B determination by single radial immunodiffusion and immunonephelometric assay (42). As we have shown for such LDL samples, where the presence of aggregates was confirmed by non-denaturing gradient polyacrylamide gel electrophoresis with subsequent immunoblotting, the apparent loss of apolipoprotein B immunoreactivity (up to 23%) is observed only when "gel" immunochemical techniques (single radial immunodiffusion and electroimmunoassay) are used in contrast to immunoturbidimetric assay.

It is obvious that aggregation of LDL can lead to the alterations not only in immunochemical, but other (physico-chemical, ability to bind with B,E-receptors and etc.) native properties of LDL. The electrophoretic and other data obtained by us show that the degree of aggregation of LDL during low-temperature storage is not dependent as much upon the duration of their storage as directly upon the freeze-thawing process, and dramatically increases (even up to opalescence of the solution and visible precipitation) at multi-fold repetition of the operation. The addition of 1.4 mol/l dimethyl sulphoxide prior to freezing of LDL and their subsequent long-term storage at low temperature prevents such LDL aggregation and allows them to maintain their native properties. This conclusion was also confirmed by very recently published data (12), where the authors used sucrose for cryopreservation of LDL. The utilization of bovine serum albumin, HDL, and apolipoprotein A-I, having certain antiaggregatory effect with regard to LDL, to stabilize LDL (38, 40) is impossible in many fields of LDL application (for example, by examination of physical and chemical properties of LDL, by obtaining antibodies against apolipoprotein B, etc.). Thus, we have concentrated our efforts on the search for a relevant "cryoprotector", exercising minimal influence upon native LDL properties and allowing alterations in these properties to

be avoided during the freeze-thawing process. Among cell cryoprotectors used to preserve such complex biological structures as cells (13, 14), we have selected two of the most widely used, i.e. glycerol and dimethyl sulphoxide. Nevertheless, the results of the preliminary experiments on the utilization of glycerol as a cryoprotector to stabilize LDL were negative, leading us to further study the possibility of application of only dimethyl sulphoxide to preserve LDL.

The data obtained show that the addition of 1.4 mol/l dimethyl sulphoxide directly to the solution of freshly isolated LDL in its high salt buffer and subsequent freezing in accordance with the above method allow LDL to be stored at -70 °C (or preferably in liquid nitrogen) for a considerable period (for up to at least 2 years, and perhaps even longer) without any alterations in their native properties. Moreover, as we also used highly concentrated LDL samples (10–30 g/l by protein), a number of duplicated experiments performed have shown that in many cases (immunochemical apolipoprotein B determination, LDL binding with B,E-receptors, animal immunization, electrophoretic methods of analysis, electron microscopy, etc.), when LDL are used in a highly diluted state, preliminary dialysis of the "preserved" LDL samples is not necessary. Using [¹²⁵I]LDL as an example, we have shown the possibility of storage of radiolabelled LDL samples, but in this case the storage time is probably limited by the lifetimes of the relevant radioisotopes. Our observations also strongly suggest that the thermostability of LDL in the presence of 1.4 mol/l dimethyl sulphoxide is rather higher (*A. B. Sigalov*, unpublished data). The absence of significant differences in the properties of the freshly isolated LDL and the LDL preparations cryopreserved according to our methods, as well as the stability of these properties during long-term storage, prove the minimal impact of dimethyl sulphoxide upon the structure and properties of lipoprotein particles, although its destabilizing effect on the secondary protein structure was reported (43).

Thus, the data obtained indicate that dimethyl sulphoxide can be successfully used as a cryoprotector for long-term storage of LDL at low temperature as well as sucrose (12) at least in the applications used. However, it is quite possible that in any particular applications one of them will be preferable. The given methods of LDL cryopreservation may be quite universal with regard to other classes of human lipoproteins and at present we are conducting relevant studies in this direction.

In conclusion we would like to express our hope that the suggested simple and effective technique of LDL storage might find wide application in clinical and biomedical investigations.

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